10/594908

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SEOUENCE LISTING

```
<110> Wang, Xiangbin
     Huang, Hualiang
      Zhao, Baofeng
     Zhao, Qi
     Piao, Jinhua
     Lin, Qing
<120> A GENETIC ENGINEERING RECOMBINANT ANTI-CEA, ANTI-CD3
      AND ANTI-CD28 SINGLE-CHAIN TRI-SPECIFIC ANTIBODY
<130> 11774-006-999 (I040179)
<150> PCT/CN2005/000408
<151> 2005-03-29
<150> CN 200410032158.3
<151> 2004-04-01
<160> 58
<170> FastSEQ for Windows Version 4.0
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<223> murine anti-CEA single chain fragment of variable region
      contained in CEA-scTsAb
Gln Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Met Lys Pro Gly Ala
Ser Val Lys Ile Ser Cys Lys Ala Thr Gly Tyr Thr Phe Ser Asp Tyr
                                25
Trp Ile Glu Trp Val Lys Gln Arg Pro Gly His Gly Leu Glu Trp Ile
                            40
Gly Glu Ile Leu Pro Gly Ser Gly Arg Thr Asp Tyr Asn Glu Arg Phe
                        55
Lys Gly Lys Ala Thr Phe Thr Gly Asp Val Ser Ser Asn Thr Ala Tyr
                    70
                                        75
Met Lys Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys
                85
                                    90
Ala Thr Gly Thr Thr Pro Phe Gly Tyr Trp Gly Gln Gly Thr Leu Val
            100
                                105
Thr Val Ser Ala Thr Ser Thr Pro Ser His Asn Ser His Gln Val Pro
                            120
                                                 125
Ser Ala Gly Gly Pro Thr Ala Asn Ser Gly Ser Arg Asp Ile Val Leu
                                            140
                        135
Thr Gln Ser Pro Ala Ser Leu Ala Val Ser Leu Gly Gln Arg Ala Thr
                    150
                                        155
Ile Ser Cys Arg Ala Ser Gln Ser Val Ser Thr Ser Ser Tyr Thr Tyr
                                    170
                165
Met His Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro Lys Leu Ile
            180
                                185
                                                     190
Lys Tyr Ala Ser Asn Leu Glu Ser Gly Val Pro Ala Arg Phe Ser Gly
                            200
                                                 205
        195
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Ser Gly Ser Gly Thr Asp Phe Thr Leu Asn Ile His Pro Val Glu Glu
    210
                        215
Glu Asp Thr Ala Tyr Tyr Tyr Cys Gln His Ser Trp Glu Ile Pro Arg
                    230
                                        235
Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
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<223> murine anti-CEA single chain fragment of variable region
     contained in CEA-scTsAb
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Glu Val Lys Leu Val Glu Ser Gly Pro Glu Leu Val Lys Pro Gly Ala
                                   10
Ser Met Lys Ile Ser Cys Lys Ala Ser Gly Tyr Ser Phe Thr Gly Tyr
                                25
Thr Met Asn Trp Val Lys Gln Ser His Gly Lys Asn Leu Glu Trp Met
Gly Leu Ile Asn Pro Tyr Lys Gly Val Ser Thr Tyr Asn Gln Lys Phe
Lys Asp Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser Thr Ala Tyr
Met Glu Leu Leu Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys
                                    90
Ala Arg Ser Gly Tyr Tyr Gly Asp Ser Asp Trp Tyr Phe Asp Val Trp
                               105
Gly Ala Gly Thr Ser Val Thr Val Ser Ser Thr Ser Gly Gly Gly
                           120
Ser Gly Gly Gly Ser Gly Gly Gly Ser Ser Arg Asp Ile Gln
                       135
Met Thr Gln Thr Thr Ser Ser Leu Ser Ala Ser Leu Gly Asp Arg Val
                   150
                                        155
Thr Ile Ser Cys Arg Ala Ser Gln Asp Ile Arg Asn Tyr Leu Asn Trp
                                   170
Tyr Gln Gln Lys Pro Asp Gly Thr Val Lys Leu Leu Ile Tyr Tyr Thr
                               185
Ser Arg Leu His Ser Gly Val Pro Ser Lys Phe Ser Gly Ser Gly Ser
                           200
Gly Thr Asp Tyr Ser Leu Thr Ile Ser Asn Leu Glu Gln Glu Asp Ile
                       215
                                            220
Ala Thr Tyr Phe Cys Gln Gln Gly Asn Thr Leu Pro Trp Thr Phe Ala
                   230
                                        235
Gly Gly Thr Lys Leu Glu Leu Lys Arg Ala
               245
<210> 3
<211> 2103
<212> DNA
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<220>
<223> CEA-scTsAb sequence
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<400> 3

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atgggtctcg agcaggtgca gctgcagcag agcggtgcgg aactgatgaa accgggcgcg 60
agcgtgaaaa tcagctgcaa agcgaccggc tataccttca gcgattattg gatcgaatgg 120
gtgaaacagc gtccgggtca cggcctggaa tggatcggtg aaatcctgcc gggcagcggc 180
cgtaccgact acaacgaacg tttcaaaggc aaagcgacct tcaccggcga cgtttctagc 240
aacaccgcgt atatgaaact gtctagcctg accagcgaag atagcgcggt gtattactgc 300
gcgaccggca ccaccccgtt cggttactgg ggtcagggca ccctggttac cgtttccgcg 360
actagtacco cgagocataa cagocatcag gtgccgagog cgggcggccc gaccgcgaac 420
ageggeteta gagacategt getgaeceag ageceggega geetggeggt gtetetgggt 480
cagcgtgcga ccatctcctg ccgtgcttcc cagtccgttt ccacctcctc ctacacctac 540
atgcactggt atcagcagaa accgggtcag ccgccgaaac tgctgatcaa atatgcgagc 600
aacctggaat ctggtgtgcc ggcgcgtttc agcggttctg gcagcggcac cgacttcacc 660
ctgaacatcc accoggtgga agaagaagat accgcgtatt actattgcca gcactcttgg 720
gaaatcccgc gtaccttcgg tggcggcacc aaactggaaa tcaaagaatt caacagcacg 780
taccgggttg taagcgtcct caccgtactg caccaggact ggctgaatgg caaggaatac 840
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getteaatga agatateetg caaggettet ggttaeteat teaetggeta caecatgaac 960
tgggtgaagc agagtcatgg aaagaacctt gagtggatgg gacttattaa tccttacaaa 1020
ggtgttagta cctacaacca gaagttcaag gacaaggcca cattaactgt agacaagtca 1080
tccagcacag cctacatgga actcctcagt ctgacatctg aggactctgc agtctattac 1140
tgtgcaagat cggggtacta cggtgatagt gactggtact tcgatgtctg gggcgcagga 1200
acctcagtca ctgtctcctc aactagtggt ggtggttggtt ctggtggtgg tggttctggt 1260
ggtggtggtt cttctagaga catccagatg acccagacca catcctccct gtctgcctct 1320
ctgggagaca gagtcaccat cagttgcagg gcaagtcagg acattagaaa ttatttaaac 1380
tggtatcaac agaaaccaga tggaactgtt aaactcctga tctactacac atcaagatta 1440
cactcaggag teccateaaa gtteagtgge agtgggtetg gaacagatta tteteteace 1500
attagcaacc tggagcaaga ggatattgcc acttactttt gccaacaggg taatacgctt 1560
ccgtggacgt tcgctggagg caccaaactg gaactgaagc gcgctgtcga cttccagaat 1620
gcgctgctgg ttcgttacac caagaaagta ccccaagtgt caactccaac tcctgtagag 1680
gtctcacata tgcaggtaca gctacaggaa tctggtccgg gtctggtaaa accgtctcag 1740
accetytete tyacetytae egtatetyyt ttetetetyt etyactatyy tytteattyy 1800
gtacgtcagc cgccaggtaa aggtctggaa tgtctgggtg taatatgggg tggaggcacg 1860
aattataatt eggeteteat gteeagaegt gtaacetett eegaegatae etetaaaaat 1920
cagttetete tgaaactgte tteegtagae accgetgtat actattgtge tegtteetat 1980.
tactattcta tggactactg gggtcagggc accctggtaa ccgtatcttc cggtaccgaa 2040
caaaaactca totoagaaga ggatotgaat ggggccgcac atcatcatca coatcacgag 2100
                                                                   2103
caa
<210> 4
<211> 701
<212> PRT
<213> Artificial Sequence
<220>
<223> CEA-scTsAb sequence
<400> 4
Met Gly Leu Glu Gln Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Met
Lys Pro Gly Ala Ser Val Lys Ile Ser Cys Lys Ala Thr Gly Tyr Thr
                                25
Phe Ser Asp Tyr Trp Ile Glu Trp Val Lys Gln Arg Pro Gly His Gly
                            40
Leu Glu Trp Ile Gly Glu Ile Leu Pro Gly Ser Gly Arg Thr Asp Tyr
                        55
Asn Glu Arg Phe Lys Gly Lys Ala Thr Phe Thr Gly Asp Val Ser Ser
                                        75
Asn Thr Ala Tyr Met Lys Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala
                                    90
Val Tyr Tyr Cys Ala Thr Gly Thr Thr Pro Phe Gly Tyr Trp Gly Gln
                                105
Gly Thr Leu Val Thr Val Ser Ala Thr Ser Thr Pro Ser His Asn Ser
```

His Gln Val Pro Ser Ala Gly Gly Pro Thr Ala Asn Ser Gly Ser Arg Asp Ile Val Leu Thr Gln Ser Pro Ala Ser Leu Ala Val Ser Leu Gly Gln Arg Ala Thr Ile Ser Cys Arg Ala Ser Gln Ser Val Ser Thr Ser Ser Tyr Thr Tyr Met His Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro Lys Leu Leu Ile Lys Tyr Ala Ser Asn Leu Glu Ser Gly Val Pro Ala Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Asn Ile His Pro Val Glu Glu Glu Asp Thr Ala Tyr Tyr Tyr Cys Gln His Ser Trp Glu Ile Pro Arg Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Glu Phe Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Ser Thr Glu Val Lys Leu Val Glu Ser Gly Pro Glu Leu Val Lys Pro Gly Ala Ser Met Lys Ile Ser Cys Lys Ala Ser Gly Tyr Ser Phe Thr Gly Tyr Thr Met Asn Trp Val Lys Gln Ser His Gly Lys Asn Leu Glu Trp Met Gly Leu Ile Asn Pro Tyr Lys Gly Val Ser Thr Tyr Asn Gln Lys Phe Lys Asp Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser Thr Ala Tyr Met Glu Leu Leu Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys Ala Arg Ser Gly Tyr Tyr Gly Asp Ser Asp Trp Tyr Phe Asp Val Trp Gly Ala Gly Thr Ser Val Thr Val Ser Ser Thr Ser Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Ser Ser Arg Asp Ile Gln Met Thr Gln Thr Thr Ser Ser Leu Ser Ala Ser Leu Gly Asp Arg Val Thr Ile Ser Cys Arg Ala Ser Gln Asp Ile Arg Asn Tyr Leu Asn Trp Tyr Gln Gln Lys Pro Asp Gly Thr Val Lys Leu Leu Ile Tyr Tyr Thr Ser Arg Leu His Ser Gly Val Pro Ser Lys Phe Ser Gly Ser Gly Ser Gly Thr Asp Tyr Ser Leu Thr Ile Ser Asn Leu Glu Glu Glu Asp Ile Ala Thr Tyr Phe Cys Gln Gln Gly Asn Thr Leu Pro Trp Thr Phe Ala Gly Gly Thr Lys Leu Glu Leu Lys Arg Ala Val Asp Phe Gln Asn Ala Leu Leu Val Arg Tyr Thr Lys Lys Val Pro Gln Val Ser Thr Pro Thr Pro Val Glu Val Ser His Met Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu Val Lys Pro Ser Gln Thr Leu Ser Leu Thr Cys Thr Val Ser Gly Phe Ser Leu Ser Asp Tyr Gly Val His Trp Val Arg Gln Pro Pro Gly Lys Gly

```
Leu Glu Cys Leu Gly Val Ile Trp Gly Gly Gly Thr Asn Tyr Asn Ser
    610
Ala Leu Met Ser Arg Arg Val Thr Ser Ser Asp Asp Thr Ser Lys Asn
625
                    630
                                         635
Gln Phe Ser Leu Lys Leu Ser Ser Val Asp Thr Ala Val Tyr Tyr Cys
                645
                                     650
Ala Arg Ser Tyr Tyr Tyr Ser Met Asp Tyr Trp Gly Gln Gly Thr Leu
            660
                                 665
                                                     670
Val Thr Val Ser Ser Gly Thr Glu Gln Lys Leu Ile Ser Glu Glu Asp
        675
                            680
Leu Asn Gly Ala Ala His His His His His Glu Gln
                        695
<210> 5
<211> 18
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 5
                                                                    18
tataccatgg gtctcgag
<210> 6
<211> 59
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 6
tataccatgg gtctcgagat gtacccgcgc ggtaacacta gtgaattcaa cagcacgta 59
<210> 7
<211> 59
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 7
agccagtcct ggtgcagtac ggtgaggacg cttacaaccc ggtacgtgct gttgaattc 59
<210> 8
<211> 59
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 8
ctgcaccagg actggctgaa tggcaaggaa tacaaatgca agagtacttc tagaatgta 59
```

```
<210> 9
<211> 59
<212> DNA
<213> Artificial Sequence
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 9
cgaaccagca gcgcattctg gaagtcgacg ttaccgcgcg ggtacattct agaagtact 59
<210> 10
<211> 59
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 10
aatgegetge tggttegtta caccaagaaa gtaccccaag tgtcaactee aacteetgt 59
<210> 11
<211> 59
<212> DNA
<213> Artificial Sequence
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 11
geggtacegt tacegegegg gtacateata tgtgagacet ctacaggagt tggagttga 59
<210> 12
<211> 59
<212> DNA
<213> Artificial Sequence
<223> Synthetic fragment for making construct of
      CEA-scTsAb
cgcggtaacg gtaccgcgct ggaagttgac gaaacctacg ttccgaaaga atttaacgc 59
<210> 13
<211> 64
<212> DNA
<213> Artificial Sequence
<223> Synthetic fragment for making construct of
      CEA-scTsAb
tcgctagccc catccgcggg atgtcagcgt ggaaggtgaa ggtttccgcg ttaaattctt 60
tcgg
```

```
<210> 14
<211> 59
<212> DNA
<213> Artificial Sequence
<223> Synthetic fragment for making construct of
     CEA-scTsAb
<400> 14
atcgagetea tgtaccegeg eggtaaeget agegaacaaa aacteatete agaagagga 59
<210> 15
<211> 59
<212> DNA
<213> Artificial Sequence
<223> Synthetic fragment for making construct of
     CEA-scTsAb
tattgctcgt gatggtgatg atgatgtgcg gccccattca gatcctcttc tgagatgag 59
<210> 16
<211> 30
<212> DNA
<213> Artificial Sequence
<223> Synthetic fragment for making construct of
     CEA-scTsAb
<400> 16
                                                                    30
ctcgacggat ccttattgct cgtgatggtg
<210> 17
<211> 22
<212> DNA
<213> Artificial Sequence
<223> Synthetic fragment for making construct of
     CEA-scTsAb
<400> 17
                                                                    22
taatacgact cactataggg ga
<210> 18
<211> 19
<212> DNA
<213> Artificial Sequence
<223> Synthetic fragment for making construct of
     CEA-scTsAb
<400> 18
                                                                    19
gctagttatt gctcagcgg
```

```
<210> 19
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 19
tcacatatgc aggtacagct acag
                                                                   24
<210> 20
<211> 25
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 20
ttcgctagcg gaagatacgg tacca
                                                                   25
<210> 21
<211> 25
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 21
aagagtactg aggtgaagct ggtgg
                                                                   25
<210> 22
<211> 27
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 22
gaagtcgaca gcgcgcttca gttccag
                                                                   27
<210> 23
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> Synthetic linker peptide linking VL and VH of
      anti-CEA monoclonal antibody
<400> 23
Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser
                                    10
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<210> 24
<211> 20
<212> DNA
<213> Artificial Sequence
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 24
                                                                    20
ttcctcgagc aggttcagct
<210> 25
<211> 58
<212> DNA
<213> Artificial Sequence
<223> Synthetic fragment for making construct of
      CEA-scTsAb
                                                                    58
tegegeeegg ttteateagt teegeaeege tetgetgeag etgaaeetge tegaggaa
<210> 26
<211> 58
<212> DNA
<213> Artificial Sequence
<223> Synthetic fragment for making construct of
      CEA-scTsAb
                                                                    58
actgatgaaa ccgggcgcga gcgtgaaaat cagctgcaaa gcgaccggct ataccttc
<210> 27
<211> 44
<212> DNA
<213> Artificial Sequence
<223> Synthetic fragment for making construct of
      CEA-scTsAb
                                                                    44
cacccattcg atccaataat cgctgaaggt atagccggtc gctt
<210> 28
<211> 59
<212> DNA
<213> Artificial Sequence
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 28
attattggat cgaatgggtg aaacagcgtc cgggtcacgg cctggaatgg atcggtgaa 59
```

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<210> 29
<211> 58
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 29
acgttcgttg tagtcggtac ggccgctgcc cggcaggatt tcaccgatcc attccagg
<210> 30
<211> 60
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 30
cgtaccgact acaacgaacg tttcaaaggc aaagcgacct tcaccggcga cgtttctagc 60
<210> 31
<211> 60
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 31
ttcgctggtc aggctagaca gtttcatata cgcggtgttg ctagaaacgt cgccggtgaa 60
<210> 32
<211> 59
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 32
tgtctagcct gaccagcgaa gatagcgcgg tgtattactg cgcgaccggc accaccccg 59
<210> 33
<211> 60
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 33
gctcacggtc accagggtgc cctgacccca gtaaccgaac ggggtggtgc cggtcgcgca 60
```

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<210> 34
<211> 59
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 34
gcaccctggt gaccgtgagc gcgactagta ccccgagcca taacagccat caggtgccg 59
<210> 35
<211> 59
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 35
gtctctagag ccgctgttcg cggtcgggcc gcccgcgctc ggcacctgat ggctgttat 59
<210> 36
<211> 58
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 36
cgaacagcqg ctctagagac atcgtqctga cccaqagccc qqcqaqcctq qcqqtqtc
                                                                    58
<210> 37
<211> 60
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 37
ctgggaagca cggcaggaga tggtcgcacg ctgacccaga gacaccgcca ggctcgccgg 60
<210> 38
<211> 59
<212> DNA
<213> Artificial Sequence
<223> Synthetic fragment for making construct of
     CEA-scTsAb
<400> 38
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totoctgccg tgcttcccag tccgtttcca cctcctccta cacctacatg cactggtat 59
<210> 39
<211> 56
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 39
                                                                   56
gatcagcagt ttcggcggct gacccggttt ctgctgatac cagtgcatgt aggtgt
<210> 40
<211> 59
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 40
agccgccgaa actgctgatc aaatatgcga gcaacctgga atctggtgtg ccggcgcgt 59
<210> 41
<211> 59
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 41
gttcagggtg aagtcggtgc cgctgccaga accgctgaaa cgcgccggca caccagatt 59
<210> 42
<211> 59
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 42
gcaccgactt caccctgaac atccacccgg tggaagaaga agataccgcg tattactat 59
<210> 43
<211> 59
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
     CEA-scTsAb
<400> 43
gccaccgaag gtacgcggga tttcccaaga gtgctggcaa tagtaatacg cggtatctt 59
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<210> 44
<211> 50
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 44
tcccgcgtac cttcggtggc ggcaccaaac tggaaatcaa agaattcgcc
                                                                     50
<210> 45
<211> 21
<212> DNA
<213> Artificial Sequence
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 45
ggcgaattct ttgatttcca g
                                                                     21
<210> 46
<211> 21
<212> DNA
<213> Artificial Sequence
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 46
ggcgaattct ttgatttcca g
                                                                     21
<210> 47
<211> 20
<212> DNA
<213> Artificial Sequence
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 47
agccgccgaa actgctgatc
                                                                     20
<210> 48
<211> 20
<212> DNA
<213> Artificial Sequence
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 48
gatcagcagt ttcggcggct
                                                                     20
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<210> 49
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 49
cgaacagcgg ctctagagac
                                                                    20
<210> 50
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 50
gtctctagag ccgctgttcg
                                                                     20
<210> 51
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
     · CEA-scTsAb
<400> 51
                                                                     20
gtaccgacta caacgaacgt
<210> 52
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic fragment for making construct of
      CEA-scTsAb
<400> 52
                                                                     20
acgttcgttg tagtcggtac
<210> 53
<211> 26
<212> PRT
<213> Artificial Sequence
<220>
<223> FC linker
<400> 53
Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp
Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys
```

20 25

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<210> 54
<211> 25
<212> PRT
<213> Artificial Sequence
<220>
<223> HSA linker 1
<400> 54
Phe Gln Asn Ala Leu Leu Val Arg Tyr Thr Lys Lys Val Pro Gln Val
1 5
Ser Thr Pro Thr Pro Val Glu Val Ser
           20
<210> 55
<211> 24
<212> PRT
<213> Artificial Sequence
<220>
<223> SHA linker 2
<400> 55
Ala Leu Glu Val Asp Glu Thr Tyr Val Pro Lys Glu Phe Asn Ala Glu
                                   10
Thr Phe Thr Phe His Ala Asp Ile
           20
<210> 56
<211> 11
<212> PRT
<213> Artificial Sequence
<220>
<223> cmyc tag
<400> 56
Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn
           5
<210> 57
<211> 6
<212> PRT
<213> Artificial Sequence
<220>
<223> His tag
<400> 57
His His His His His
<210> 58
<211> 437
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<212> DNA
<213> Artificial Sequence

<220>
<223> nucleotide sequence showing multiple cloning sites (fig 3)

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